

# Surrogate Bayesian Networks for Approximating Evolutionary Games

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## Background

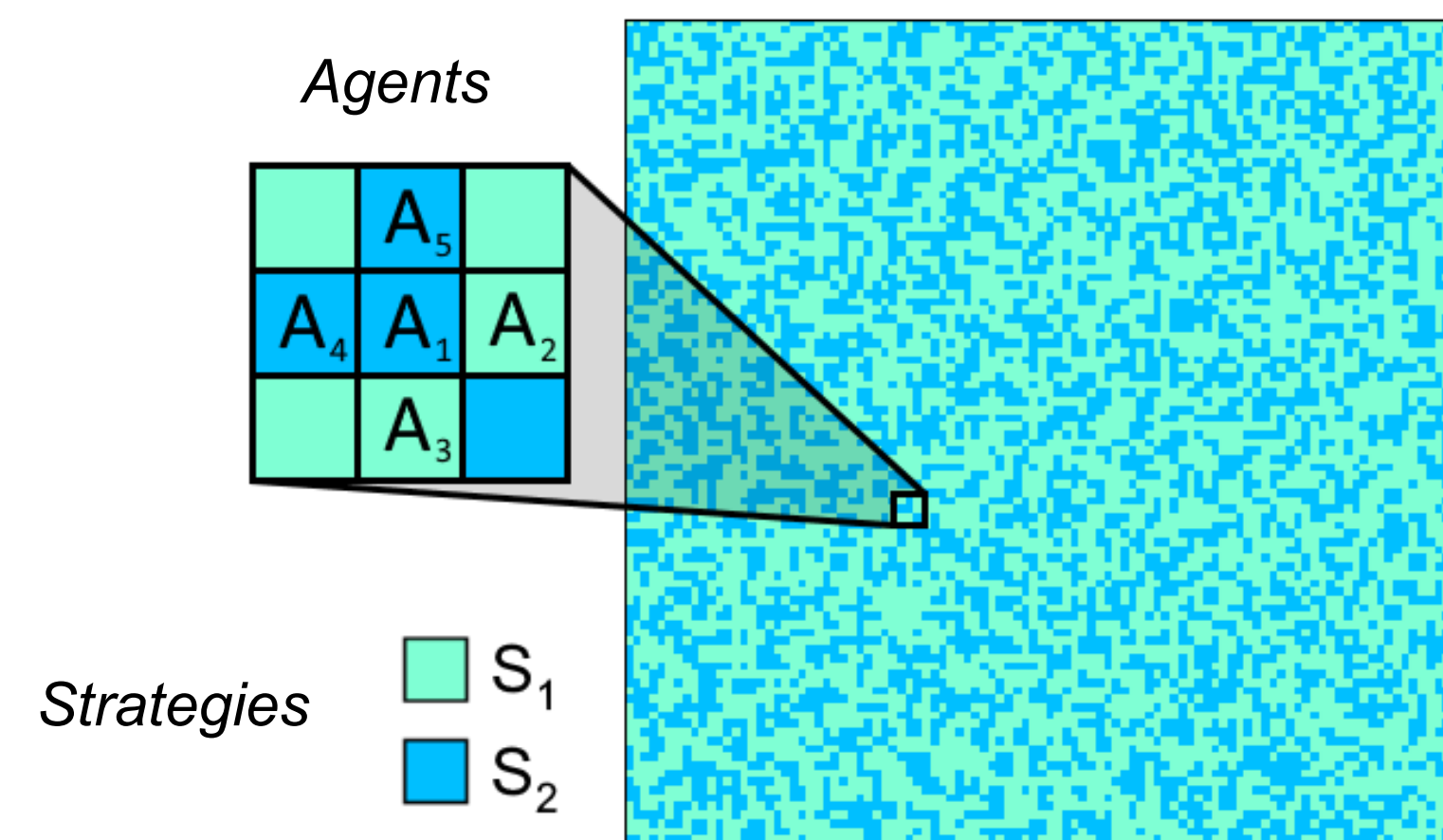
### Evolutionary Game Theory (EGT)

- Application of game theory to evolving populations

### Spatial Evolutionary Game

- EGT model on structured population (e.g. grid)
- Spatial EGT = (S, U, G, F)
  - S - set of strategies
  - U - payoff matrix/function
  - G - graph of population structure
  - F - replicator rule (e.g. Fermi rule)

$$U = \begin{matrix} & s_1 & s_2 \\ s_1 & a & b \\ s_2 & c & d \end{matrix}$$



**Goal:** evaluate population dynamics over time, e.g. proportion of pairs of adjacent agents playing strategy i and j at a given time T.

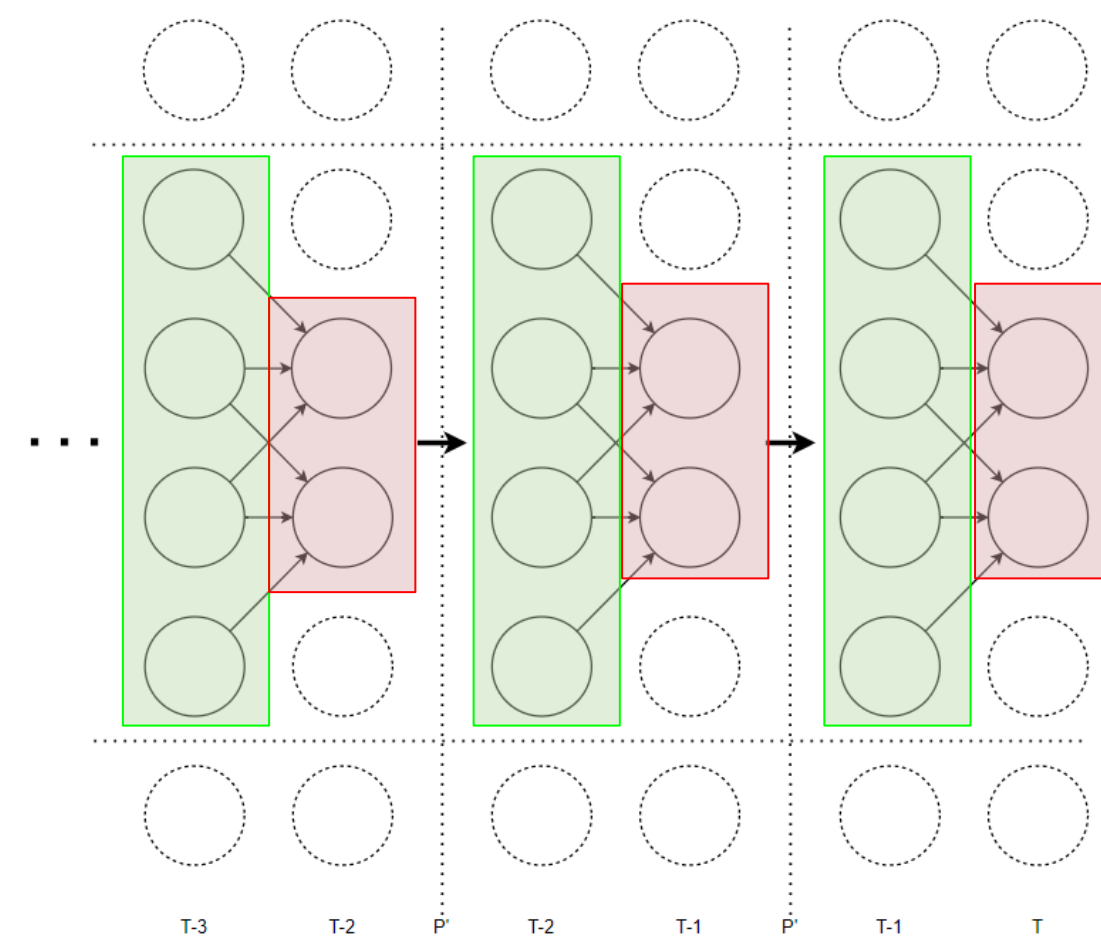
### Bayesian Network (BN)

- Bayesian Network: graphical model (X,D,F)
  - Variables:  $X = \{X_1, X_2, \dots, X_N\}$
  - Domains:  $D = \{D_{X_1}, D_{X_2}, \dots, D_{X_N}\}$
  - Parent Functions:  $F = \{F_1, F_2, \dots, F_N\}$

## Problem Statement

### Current Approach

- Use Symmetric Dynamic Bayesian Network Approximation (SD-BNA) to model spatial EGT dynamics
  - SD-BNA's have input (green) and output (red) sections at each iteration
  - Larger input  $\rightarrow$  more accurate approximation



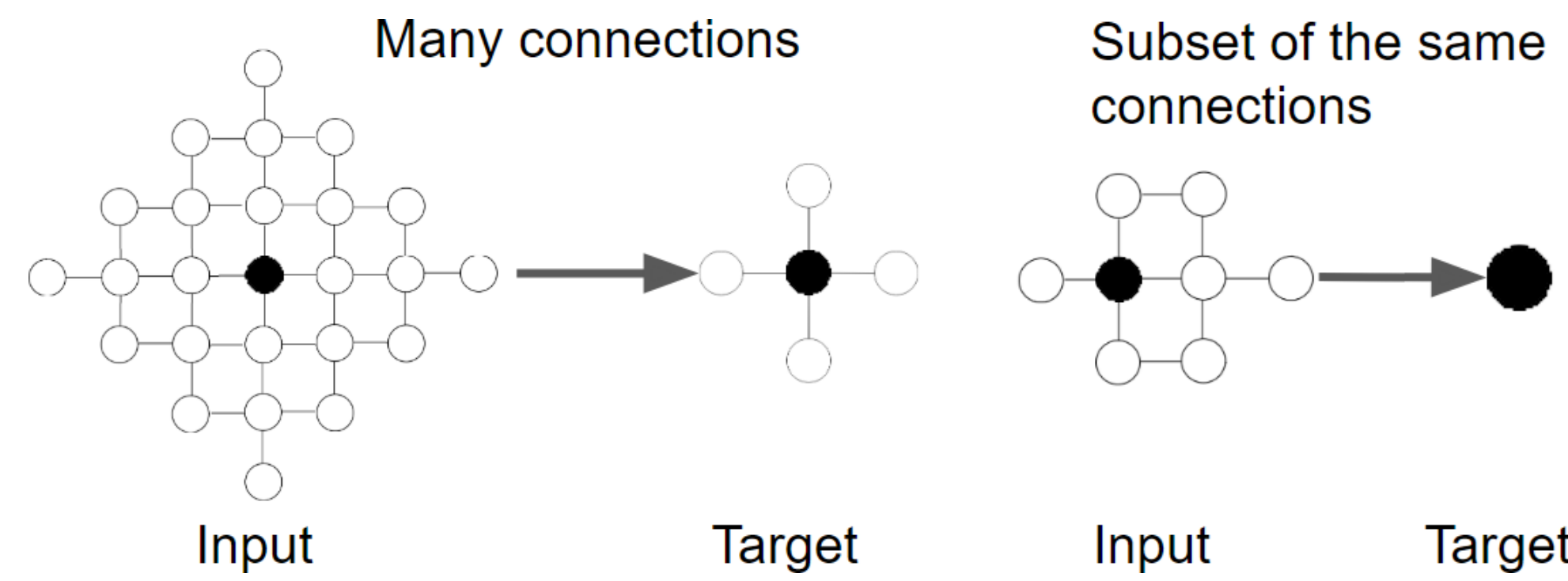
- But larger input can be computationally expensive

### Proposed Approach

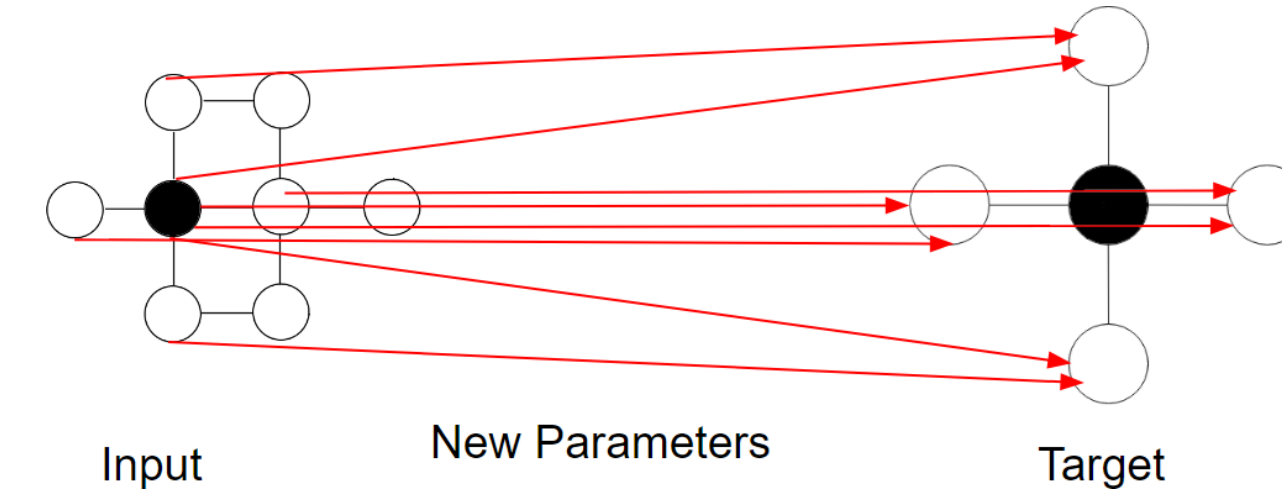
- Approximate inference on large SD-BNA, using a smaller **surrogate** SD-BNA
- Extend smaller SD-BNA with new parameters learned using samples from the large SD-BNA

## Surrogate Maximum Likelihood Models

- Consider a 25 node input SD-BNA (B) and an 8 node input SD-BNA (A)
- Smaller network is a subset of the larger network



- Add 4 dummy nodes in the output neighborhood



- New edges from input the target
- Each new node is conditioned on two nodes in the input, CPTs of the form  $\theta_{x,y,z} = P_{x|yz}$
- To learn these CPTs, we use MLE with samples from larger SD-BNA

### Basic Maximum Likelihood Estimation (MLE) algorithm

- Forward sample larger model for full configuration of every variable in the network
- The 8 node network is fully observed for each data point since its variables are subsumed in the large network
- We add symmetry constraints using pre-computed quantities from non-extended small network.
- Final problem is to maximize the following log-likelihood expression:

$$\theta^* = \arg \max_{\theta} \sum_{j=1}^L \log \theta(Nb(X^{(j)})_{t+1}, X_t^{(j)}, Nb(X^{(j)})_t)$$

subject to

$$p_{s_i}(t+1) = \sum_{s_j, s_k \in S} \theta_{s_i, s_j, s_k} \cdot p_{s_j s_k}(t), \quad \forall s_i \in S$$

### KL-Search Minimization

- Instead of random samples, combine search with sampling to get samples that reduce KL-divergence
- Search OR-search tree of large network using best-first heuristic:

$$h_{kl}(X_1=0) = [\log(P_A(X_1=0)) - \log(P_B(X_1=0))] \cdot P_B(X_1=0)$$

- $P_A(X_1=0)$  difficult to compute, approximate with Weight Mini-bucket Elimination with small i-bound (KL-Search) or single sample Monte Carlo estimate (Fast KL-Search)

## KL-Search

### Algorithm 1: KL-Search Minimization

**Input:** Two Bayesian networks: a large network  $A$ , a smaller network  $B$ , and a parameterized extended network  $B_{\theta}$  such that all nodes in  $B$  are in  $A$  ( $B \subset A$ ), a variable ordering  $o$  over  $A$ , initial distribution  $p_{yz}(t)$ , and pre-computed output distribution  $p_x(t+1)$

**Parameters:** Number of samples  $L$

**Output:**  $\theta$ , estimated parameters that minimize difference between  $A$  and  $B_{\theta}$

```

T ← the OR-search tree on A using ordering o;
OPEN ← {root(T), 0};
// frontier nodes are ordered by the 2nd value
for i = 1 → L do
  v ← OPEN.dequeue(); // remove the node
  of highest priority
  for u ∈ children(v) do
    h_kl(u) ← [log(P_A(u)) - log(P_B(u))] · P_B(u);
    Append <u, h_kl(u)> to OPEN;
  end
end
Let X be an empty list;
for v ∈ OPEN do // leaf nodes
  Forward sample x, a full configuration of A
  conditioned on the partial configuration
  represented by v;
  Append x to X;
end
Solve θ* = arg max_θ ∑_{j=1}^L log P_{B_θ}(x^j) · P_A(x^j),
subject to
p_{s_i}(t+1) = ∑_{s_j, s_k ∈ S} θ_{s_i, s_j, s_k} · p_{s_j s_k}(t), ∀ s_i ∈ S;
Return θ*;
    
```

### Convergence of KL-Search

**Theorem 6.0.1** (Asymptotic Convergence of KL-Search Minimization). *Let  $\theta_L$  be the result of KL-Search Minimization [Algorithm 1] given  $L$  samples. Then given a family of extended networks  $B_{\theta}$  parameterized by  $\theta$ :*

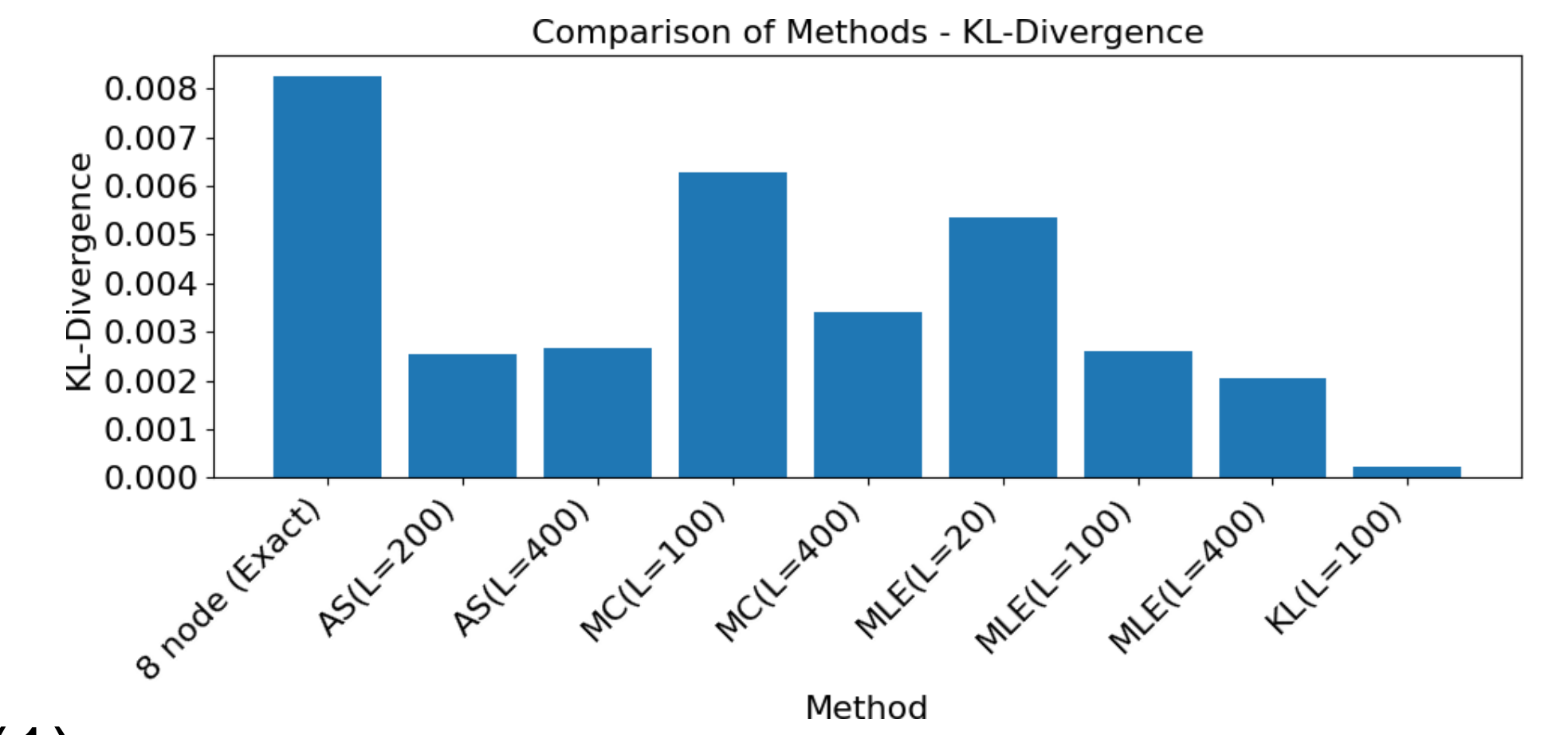
$$\lim_{L \rightarrow \infty} \theta_L = \arg \min_{\theta} D_{KL}(P_A || P_{B_{\theta}})$$

### Experimental Setup

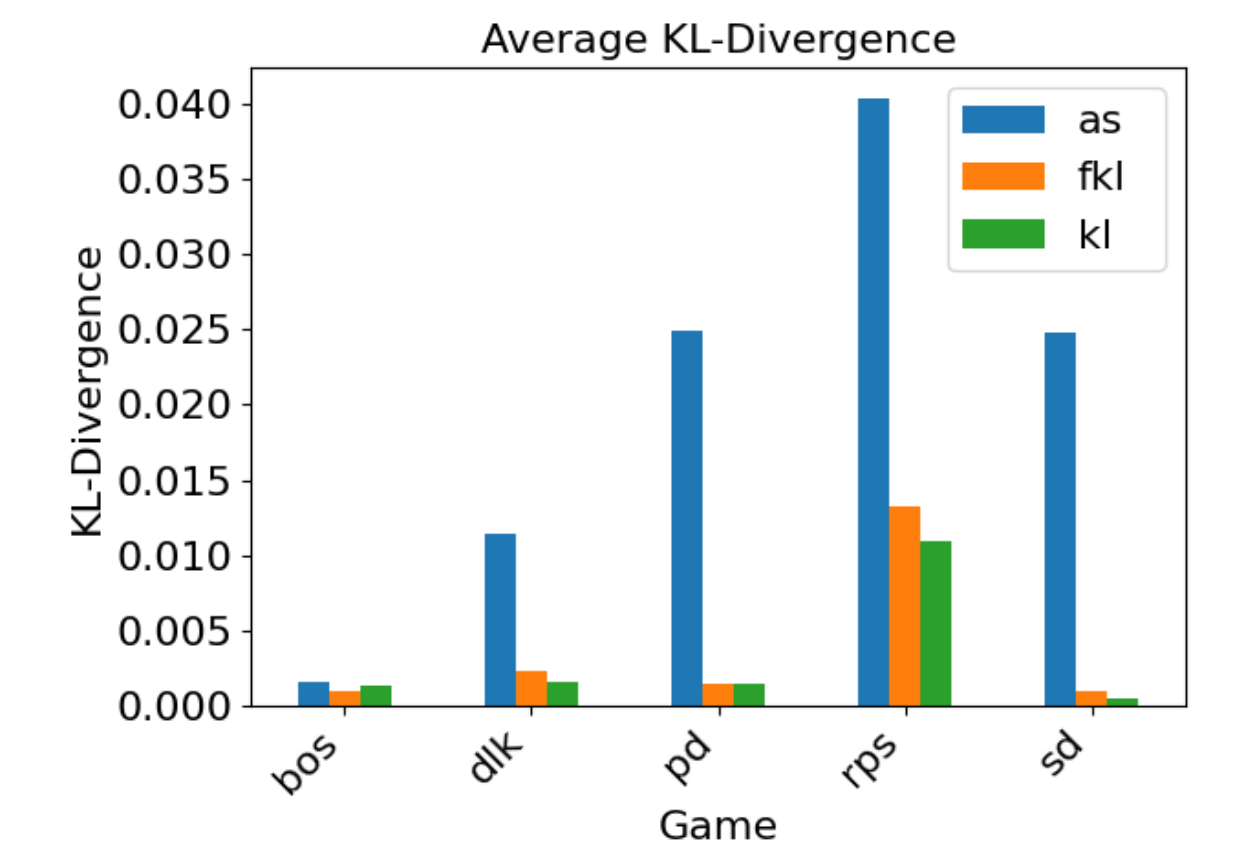
- Compare with existing approaches such as Pair Approximation and Abstraction Sampling (AS)
- Task: estimate joint probability distribution of nodes in output distribution (e.g. nodes in red area at each iteration)
- Use KL-divergence between estimate generated by simulation (ground truth) and estimate from methods begin compared.

$$D_{KL}(P_{sim}(X_1 | Nb(X)_1) || P_{method}(X_1 | Nb(X)_1))$$

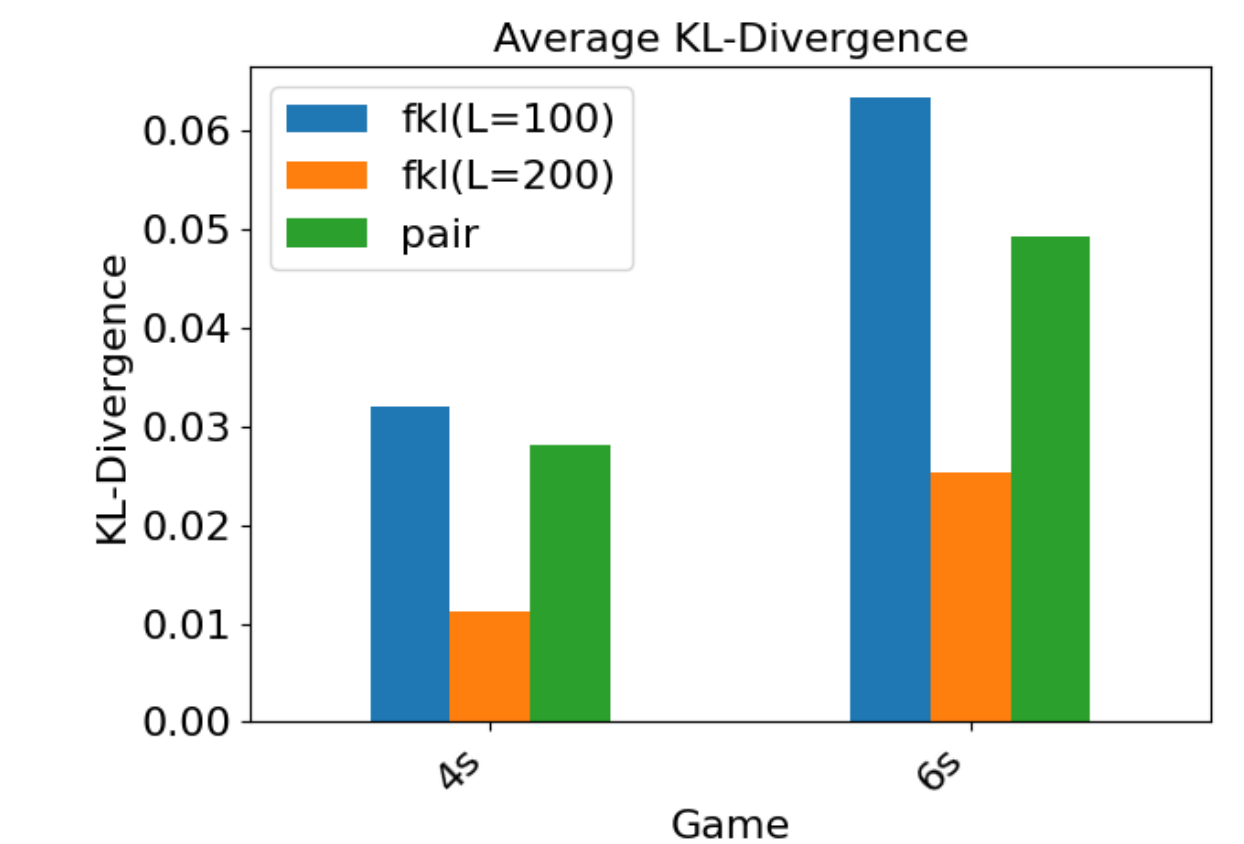
## Results



(1)



(2)



(3)

- KL-Search outperforms other methods by a considerable amount on the Deadlock game
- KL-Search and Fast KL-Search outperform Abstraction Sampling in several 2 and 3 strategy evolutionary games
- Fast KL-Search outperforms pair approximation on games with more than 4 strategies when given more than 100 samples

### Time per sample/probe

S	AS	KL-Search	Fast KL-Search
2	0.1175	0.01419	0.01624
3	0.2203	0.03063	0.03725
4	-	-	0.1630
6	-	-	0.7299

Fast KL-Search can be applied to high strategy games since it does not need to compute WMBE.

### Future Research

- Apply to domains with high degree of symmetry beyond spatial evolutionary games

### Acknowledgements

This work supported in part by NSF grant IIS-2008516 and AFOSR grant 1010GWA357.